

## SEQUENCE LISTING

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<120> GSK3 POLYPEPTIDES

<130> 59516-162/PP-15876.002/200130.524

<140> US10/211,412

<141> 2002-07-31

<150> US09/916,109

<151> 2001-07-25

<160> 11

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 420

<212> PRT

<213> Homo sapiens

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			20					25					30		
Asp	Gly	Ser	Lys	Val	Thr	Thr	Val	Val	Ala	Thr	Pro	Gly	Gln	Gly	Pro
		35					40					45			
Asp	Arg	Pro	Gln	Glu	Val	Ser	Tyr	Thr	Asp	Thr	Lys	Val	Ile	Gly	Asn
	50					55					60				
Gly	Ser	Phe	Gly	Val	Val	Tyr	Gln	Ala	Lys	Leu	Cys	Asp	Ser	Gly	Glu
65				70						75					80
Leu	Val	Ala	Ile	Lys	Lys	Val	Leu	Gln	Asp	Lys	Arg	Phe	Lys	Asn	Arg
				85					90					95	
Glu	Leu	Gln	Ile	Met	Arg	Lys	Leu	Asp	His	Cys	Asn	Ile	Val	Arg	Leu
			100					105						110	
Arg	Tyr	Phe	Phe	Tyr	Ser	Ser	Gly	Glu	Lys	Lys	Asp	Glu	Val	Tyr	Leu
		115					120					125			
Asn	Leu	Val	Leu	Asp	Tyr	Val	Pro	Glu	Thr	Val	Tyr	Arg	Val	Ala	Arg
		130				135					140				
His	Tyr	Ser	Arg	Ala	Lys	Gln	Thr	Leu	Pro	Val	Ile	Tyr	Val	Lys	Leu
145					150					155					160
Tyr	Met	Tyr	Gln	Leu	Phe	Arg	Ser	Leu	Ala	Tyr	Ile	His	Ser	Phe	Gly
				165				170						175	
Ile	Cys	His	Arg	Asp	Ile	Lys	Pro	Gln	Asn	Leu	Leu	Leu	Asp	Pro	Asp



Val	Pro	Glu	Thr	Val	Tyr	Arg	Val	Ala	Arg	His	Tyr	Ser	Arg	Ala	Lys
145					150					155					160
Gln	Thr	Leu	Pro	Val	Ile	Tyr	Val	Lys	Leu	Tyr	Met	Tyr	Gln	Leu	Phe
				165					170						175
Arg	Ser	Leu	Ala	Tyr	Ile	His	Ser	Phe	Gly	Ile	Cys	His	Arg	Asp	Ile
			180					185					190		
Lys	Pro	Gln	Asn	Leu	Leu	Leu	Asp	Pro	Asp	Thr	Ala	Val	Leu	Lys	Leu
		195					200					205			
Cys	Asp	Phe	Gly	Ser	Ala	Lys	Gln	Leu	Val	Arg	Gly	Glu	Pro	Asn	Val
210						215					220				
Ser	Tyr	Ile	Cys	Ser	Arg	Tyr	Tyr	Arg	Ala	Pro	Glu	Leu	Ile	Phe	Gly
225					230					235					240
Ala	Thr	Asp	Tyr	Thr	Ser	Ser	Ile	Asp	Val	Trp	Ser	Ala	Gly	Cys	Val
				245					250						255
Leu	Ala	Glu	Leu	Leu	Leu	Gly	Gln	Pro	Ile	Phe	Pro	Gly	Asp	Ser	Gly
			260					265					270		
Val	Asp	Gln	Leu	Val	Glu	Ile	Ile	Lys	Val	Leu	Gly	Thr	Pro	Thr	Arg
		275					280					285			
Glu	Gln	Ile	Arg	Glu	Met	Asn	Pro	Asn	Tyr	Thr	Glu	Phe	Lys	Phe	Pro
290						295					300				
Gln	Ile	Lys	Ala	His	Pro	Trp	Thr	Lys	Val	Phe	Arg	Pro	Arg	Thr	Pro
305					310					315					320
Pro	Glu	Ala	Ile	Ala	Leu	Cys	Ser	Arg	Leu	Leu	Glu	Tyr	Thr	Pro	Thr
				325					330					335	
Ala	Arg	Leu	Thr	Pro	Leu	Glu	Ala	Cys	Ala	His	Ser	Phe	Phe	Asp	Glu
			340					345					350		
Leu	Arg	Asp	Pro	Asn	Val	Lys	His	Pro	Asn	Gly	Arg	Asp	Thr	Pro	Ala
		355					360					365			
Leu	Phe	Asn	Phe	Thr	Thr	Gln	Glu	Leu	Ser	Ser	Asn	Pro	Pro	Leu	Ala
370						375					380				
Thr	Ile	Leu	Ile	Pro	Pro	His	Ala	Arg	Ile						
385					390										

<210> 3  
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 <212> PRT  
 <213> Homo sapiens

Met	Glu	Tyr	Met	Pro	Met	Glu	Gly	Gly	Gly	Gly	Ser	Lys	Val	Thr	Thr
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			20					25					30		
Tyr	Thr	Asp	Thr	Lys	Val	Ile	Gly	Asn	Gly	Ser	Phe	Gly	Val	Val	Tyr
		35					40					45			
Gln	Ala	Lys	Leu	Cys	Asp	Ser	Gly	Glu	Leu	Val	Ala	Ile	Lys	Lys	Val
50						55					60				
Leu	Gln	Asp	Lys	Arg	Phe	Lys	Asn	Arg	Glu	Leu	Gln	Ile	Met	Arg	Lys
65					70					75					80
Leu	Asp	His	Cys	Asn	Ile	Val	Arg	Leu	Arg	Tyr	Phe	Phe	Tyr	Ser	Ser
				85					90					95	
Gly	Glu	Lys	Lys	Asp	Glu	Val	Tyr	Leu	Asn	Leu	Val	Leu	Asp	Tyr	Val
			100					105					110		
Pro	Glu	Thr	Val	Tyr	Arg	Val	Ala	Arg	His	Tyr	Ser	Arg	Ala	Lys	Gln
		115					120					125			
Thr	Leu	Pro	Val	Ile	Tyr	Val	Lys	Leu	Tyr	Met	Tyr	Gln	Leu	Phe	Arg

130		135		140
Ser Leu Ala Tyr Ile His	Ser Phe Gly Ile Cys His Arg Asp Ile Lys			
145	150	155		160
Pro Gln Asn Leu Leu Leu Asp Pro Asp Thr Ala Val Leu Lys Leu Cys				
	165	170		175
Asp Phe Gly Ser Ala Lys Gln Leu Val Arg Gly Glu Pro Asn Val Ser				
	180	185		190
Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala Pro Glu Leu Ile Phe Gly Ala				
	195	200		205
Thr Asp Tyr Thr Ser Ser Ile Asp Val Trp Ser Ala Gly Cys Val Leu				
	210	215		220
Ala Glu Leu Leu Leu Gly Gln Pro Ile Phe Pro Gly Asp Ser Gly Val				
225	230	235		240
Asp Gln Leu Val Glu Ile Ile Lys Val Leu Gly Thr Pro Thr Arg Glu				
	245	250		255
Gln Ile Arg Glu Met Asn Pro Asn Tyr Thr Glu Phe Lys Phe Pro Gln				
	260	265		270
Ile Lys Ala His Pro Trp Thr Lys Val Phe Arg Pro Arg Thr Pro Pro				
	275	280		285
Glu Ala Ile Ala Leu Cys Ser Arg Leu Leu Glu Tyr Thr Pro Thr Ala				
	290	295		300
Arg Leu Thr Pro Leu Glu Ala Cys Ala His Ser Phe Phe Asp Glu Leu				
305	310	315		320
Arg Asp Pro Asn Val Lys His Pro Asn Gly Arg Asp Thr Pro Ala Leu				
	325	330		335
Phe Asn Phe Thr Thr Gln Glu Leu Ser Ser Asn Pro Pro Leu Ala Thr				
	340	345		350
Ile Leu Ile Pro Pro His Ala Arg Ile				
	355	360		

<210> 4

<211> 483

<212> PRT

<213> Homo sapiens

<400> 4

Met Ser Gly Gly Gly Pro Ser Gly Gly Gly Pro Gly Gly Ser Gly Arg				
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	20	25	30	
Gly Gly Gly Pro Gly Gly Ser Ala Ser Gly Pro Gly Gly Thr Gly Gly				
	35	40	45	
Gly Lys Ala Ser Val Gly Ala Met Gly Gly Gly Val Gly Ala Ser Ser				
	50	55	60	
Ser Gly Gly Gly Pro Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Pro				
65	70	75	80	
Gly Ala Gly Thr Ser Phe Pro Pro Pro Gly Val Lys Leu Gly Arg Asp				
	85	90	95	
Ser Gly Lys Val Thr Thr Val Val Ala Thr Leu Gly Gln Gly Pro Glu				
	100	105	110	
Arg Ser Gln Glu Val Ala Tyr Thr Asp Ile Lys Val Ile Gly Asn Gly				
	115	120	125	
Ser Phe Gly Val Val Tyr Gln Ala Arg Leu Ala Glu Thr Arg Glu Leu				
	130	135	140	
Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Phe Lys Asn Arg Glu				
145	150	155	160	

Leu	Gln	Ile	Met	Arg	Lys	Leu	Asp	His	Cys	Asn	Ile	Val	Arg	Leu	Arg
			165						170					175	
Tyr	Phe	Phe	Tyr	Ser	Ser	Gly	Glu	Lys	Lys	Asp	Glu	Leu	Tyr	Leu	Asn
			180					185					190		
Leu	Val	Leu	Glu	Tyr	Val	Pro	Glu	Thr	Val	Tyr	Arg	Val	Ala	Arg	His
		195					200					205			
Phe	Thr	Lys	Ala	Lys	Leu	Thr	Ile	Pro	Ile	Leu	Tyr	Val	Lys	Val	Tyr
	210					215					220				
Met	Tyr	Gln	Leu	Phe	Arg	Ser	Leu	Ala	Tyr	Ile	His	Ser	Gln	Gly	Val
225					230					235				240	
Cys	His	Arg	Asp	Ile	Lys	Pro	Gln	Asn	Leu	Leu	Val	Asp	Pro	Asp	Thr
			245						250					255	
Ala	Val	Leu	Lys	Leu	Cys	Asp	Phe	Gly	Ser	Ala	Lys	Gln	Leu	Val	Arg
		260						265					270		
Gly	Glu	Pro	Asn	Val	Ser	Tyr	Ile	Cys	Ser	Arg	Tyr	Tyr	Arg	Ala	Pro
	275						280					285			
Glu	Leu	Ile	Phe	Gly	Ala	Thr	Asp	Tyr	Thr	Ser	Ser	Ile	Asp	Val	Trp
	290					295					300				
Ser	Ala	Gly	Cys	Val	Leu	Ala	Glu	Leu	Leu	Leu	Gly	Gln	Pro	Ile	Phe
305					310					315				320	
Pro	Gly	Asp	Ser	Gly	Val	Asp	Gln	Leu	Val	Glu	Ile	Ile	Lys	Val	Leu
			325						330					335	
Gly	Thr	Pro	Thr	Arg	Glu	Gln	Ile	Arg	Glu	Met	Asn	Pro	Asn	Tyr	Thr
		340						345					350		
Glu	Phe	Lys	Phe	Pro	Gln	Ile	Lys	Ala	His	Pro	Trp	Thr	Lys	Val	Phe
		355					360					365			
Lys	Ser	Arg	Thr	Pro	Pro	Glu	Ala	Ile	Ala	Leu	Cys	Ser	Ser	Leu	Leu
	370					375					380				
Glu	Tyr	Thr	Pro	Ser	Ser	Arg	Leu	Ser	Pro	Leu	Glu	Ala	Cys	Ala	His
385					390					395					400
Ser	Phe	Phe	Asp	Glu	Leu	Arg	Cys	Leu	Gly	Thr	Gln	Leu	Pro	Asn	Asn
			405						410					415	
Arg	Pro	Leu	Pro	Pro	Leu	Phe	Asn	Phe	Ser	Ala	Gly	Glu	Leu	Ser	Ile
		420						425					430		
Gln	Pro	Ser	Leu	Asn	Ala	Ile	Leu	Ile	Pro	Pro	His	Leu	Arg	Ser	Pro
	435						440					445			
Ala	Gly	Thr	Thr	Thr	Leu	Thr	Pro	Ser	Ser	Gln	Ala	Leu	Thr	Glu	Thr
	450					455					460				
Pro	Thr	Ser	Ser	Asp	Trp	Gln	Ser	Thr	Asp	Ala	Thr	Pro	Thr	Leu	Thr
465					470					475					480
Asn	Ser	Ser													

<210> 5

<211> 447

<212> PRT

<213> Homo sapiens

<400> 5

Met	Ser	Gly	Gly	Gly	Pro	Ser	Gly	Gly	Gly	Pro	Gly	Gly	Ser	Gly	Arg
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Ala	Arg	Thr	Ser	Ser	Phe	Ala	Glu	Pro	Gly	Gly	Gly	Gly	Gly	Gly	Gly
		20					25						30		
Gly	Gly	Gly	Pro	Gly	Gly	Ser	Ala	Ser	Gly	Pro	Gly	Gly	Thr	Gly	Gly
	35					40						45			
Gly	Lys	Ala	Ser	Val	Gly	Ala	Met	Gly	Gly	Gly	Val	Gly	Ala	Ser	Ser

50		55		60
Ser Gly Gly Gly Pro	Gly Gly Ser Gly Gly	Gly Gly Ser Gly Gly Pro		
65	70	75	80	
Gly Ala Gly Thr Ser	Phe Pro Pro Pro Gly	Val Lys Leu Gly Arg Asp		
	85	90	95	
Ser Gly Lys Val Thr	Thr Val Val Ala Thr	Leu Gly Gln Gly Pro Glu		
	100	105	110	
Arg Ser Gln Glu Val	Ala Tyr Thr Asp Ile	Lys Val Ile Gly Asn Gly		
	115	120	125	
Ser Phe Gly Val Val	Tyr Gln Ala Arg Leu	Ala Glu Thr Arg Glu Leu		
	130	135	140	
Val Ala Ile Lys Lys	Val Leu Gln Asp Lys	Arg Phe Lys Asn Arg Glu		
145	150	155	160	
Leu Gln Ile Met Arg	Lys Leu Asp His Cys	Asn Ile Val Arg Leu Arg		
	165	170	175	
Tyr Phe Phe Tyr Ser	Ser Gly Glu Lys Lys	Asp Glu Leu Tyr Leu Asn		
	180	185	190	
Leu Val Leu Glu Tyr	Val Pro Glu Thr Val	Tyr Arg Val Ala Arg His		
	195	200	205	
Phe Thr Lys Ala Lys	Leu Thr Ile Pro Ile	Leu Tyr Val Lys Val Tyr		
	210	215	220	
Met Tyr Gln Leu Phe	Arg Ser Leu Ala Tyr	Ile His Ser Gln Gly Val		
225	230	235	240	
Cys His Arg Asp Ile	Lys Pro Gln Asn Leu	Leu Val Asp Pro Asp Thr		
	245	250	255	
Ala Val Leu Lys Leu	Cys Asp Phe Gly Ser	Ala Lys Gln Leu Val Arg		
	260	265	270	
Gly Glu Pro Asn Val	Ser Tyr Ile Cys Ser	Arg Tyr Tyr Arg Ala Pro		
	275	280	285	
Glu Leu Ile Phe Gly	Ala Thr Asp Tyr Thr	Ser Ser Ile Asp Val Trp		
	290	295	300	
Ser Ala Gly Cys Val	Leu Ala Glu Leu Leu	Leu Gly Gln Pro Ile Phe		
305	310	315	320	
Pro Gly Asp Ser Gly	Val Asp Gln Leu Val	Glu Ile Ile Lys Val Leu		
	325	330	335	
Gly Thr Pro Thr Arg	Glu Gln Ile Arg Glu	Met Asn Pro Asn Tyr Thr		
	340	345	350	
Glu Phe Lys Phe Pro	Gln Ile Lys Ala His	Pro Trp Thr Lys Val Phe		
	355	360	365	
Lys Ser Arg Thr Pro	Pro Glu Ala Ile Ala	Leu Cys Ser Ser Leu Leu		
	370	375	380	
Glu Tyr Thr Pro Ser	Ser Arg Leu Ser Pro	Leu Glu Ala Cys Ala His		
385	390	395	400	
Ser Phe Phe Asp Glu	Leu Arg Cys Leu Gly	Thr Gln Leu Pro Asn Asn		
	405	410	415	
Arg Pro Leu Pro Pro	Leu Phe Asn Phe Ser	Ala Gly Glu Leu Ser Ile		
	420	425	430	
Gln Pro Ser Leu Asn	Ala Ile Leu Ile Pro	Pro His Leu Arg Ser		
	435	440	445	

<210> 6

<211> 387

<212> PRT

<213> Homo sapiens.

<400> 6

Ser Gly Lys Val Thr Thr Val Val Ala Thr Leu Gly Gln Gly Pro Glu  
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 Arg Ser Gln Glu Val Ala Tyr Thr Asp Ile Lys Val Ile Gly Asn Gly  
 20 25 30  
 Ser Phe Gly Val Val Tyr Gln Ala Arg Leu Ala Glu Thr Arg Glu Leu  
 35 40 45  
 Val Ala Ile Lys Lys Val Leu Gln Asp Lys Arg Phe Lys Asn Arg Glu  
 50 55 60  
 Leu Gln Ile Met Arg Lys Leu Asp His Cys Asn Ile Val Arg Leu Arg  
 65 70 75 80  
 Tyr Phe Phe Tyr Ser Ser Gly Glu Lys Lys Asp Glu Leu Tyr Leu Asn  
 85 90 95  
 Leu Val Leu Glu Tyr Val Pro Glu Thr Val Tyr Arg Val Ala Arg His  
 100 105 110  
 Phe Thr Lys Ala Lys Leu Thr Ile Pro Ile Leu Tyr Val Lys Val Tyr  
 115 120 125  
 Met Tyr Gln Leu Phe Arg Ser Leu Ala Tyr Ile His Ser Gln Gly Val  
 130 135 140  
 Cys His Arg Asp Ile Lys Pro Gln Asn Leu Leu Val Asp Pro Asp Thr  
 145 150 155 160  
 Ala Val Leu Lys Leu Cys Asp Phe Gly Ser Ala Lys Gln Leu Val Arg  
 165 170 175  
 Gly Glu Pro Asn Val Ser Tyr Ile Cys Ser Arg Tyr Tyr Arg Ala Pro  
 180 185 190  
 Glu Leu Ile Phe Gly Ala Thr Asp Tyr Thr Ser Ser Ile Asp Val Trp  
 195 200 205  
 Ser Ala Gly Cys Val Leu Ala Glu Leu Leu Leu Gly Gln Pro Ile Phe  
 210 215 220  
 Pro Gly Asp Ser Gly Val Asp Gln Leu Val Glu Ile Ile Lys Val Leu  
 225 230 235 240  
 Gly Thr Pro Thr Arg Glu Gln Ile Arg Glu Met Asn Pro Asn Tyr Thr  
 245 250 255  
 Glu Phe Lys Phe Pro Gln Ile Lys Ala His Pro Trp Thr Lys Val Phe  
 260 265 270  
 Lys Ser Arg Thr Pro Pro Glu Ala Ile Ala Leu Cys Ser Ser Leu Leu  
 275 280 285  
 Glu Tyr Thr Pro Ser Ser Arg Leu Ser Pro Leu Glu Ala Cys Ala His  
 290 295 300  
 Ser Phe Phe Asp Glu Leu Arg Cys Leu Gly Thr Gln Leu Pro Asn Asn  
 305 310 315 320  
 Arg Pro Leu Pro Pro Leu Phe Asn Phe Ser Ala Gly Glu Leu Ser Ile  
 325 330 335  
 Gln Pro Ser Leu Asn Ala Ile Leu Ile Pro Pro His Leu Arg Ser Pro  
 340 345 350  
 Ala Gly Thr Thr Leu Thr Pro Ser Ser Gln Ala Leu Thr Glu Thr  
 355 360 365  
 Pro Thr Ser Ser Asp Trp Gln Ser Thr Asp Ala Thr Pro Thr Leu Thr  
 370 375 380  
 Asn Ser Ser  
 385

<210> 7

<211> 351

<212> PRT

<213> Homo sapiens

<400> 7

Ser	Gly	Lys	Val	Thr	Thr	Val	Val	Ala	Thr	Leu	Gly	Gln	Gly	Pro	Glu
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Arg	Ser	Gln	Glu	Val	Ala	Tyr	Thr	Asp	Ile	Lys	Val	Ile	Gly	Asn	Gly
		20						25					30		
Ser	Phe	Gly	Val	Val	Tyr	Gln	Ala	Arg	Leu	Ala	Glu	Thr	Arg	Glu	Leu
		35					40					45			
Val	Ala	Ile	Lys	Lys	Val	Leu	Gln	Asp	Lys	Arg	Phe	Lys	Asn	Arg	Glu
	50					55					60				
Leu	Gln	Ile	Met	Arg	Lys	Leu	Asp	His	Cys	Asn	Ile	Val	Arg	Leu	Arg
65				70					75					80	
Tyr	Phe	Phe	Tyr	Ser	Ser	Gly	Glu	Lys	Lys	Asp	Glu	Leu	Tyr	Leu	Asn
			85					90					95		
Leu	Val	Leu	Glu	Tyr	Val	Pro	Glu	Thr	Val	Tyr	Arg	Val	Ala	Arg	His
			100					105					110		
Phe	Thr	Lys	Ala	Lys	Leu	Thr	Ile	Pro	Ile	Leu	Tyr	Val	Lys	Val	Tyr
		115				120						125			
Met	Tyr	Gln	Leu	Phe	Arg	Ser	Leu	Ala	Tyr	Ile	His	Ser	Gln	Gly	Val
	130					135					140				
Cys	His	Arg	Asp	Ile	Lys	Pro	Gln	Asn	Leu	Leu	Val	Asp	Pro	Asp	Thr
145				150					155					160	
Ala	Val	Leu	Lys	Leu	Cys	Asp	Phe	Gly	Ser	Ala	Lys	Gln	Leu	Val	Arg
			165					170						175	
Gly	Glu	Pro	Asn	Val	Ser	Tyr	Ile	Cys	Ser	Arg	Tyr	Tyr	Arg	Ala	Pro
			180					185					190		
Glu	Leu	Ile	Phe	Gly	Ala	Thr	Asp	Tyr	Thr	Ser	Ser	Ile	Asp	Val	Trp
		195				200						205			
Ser	Ala	Gly	Cys	Val	Leu	Ala	Glu	Leu	Leu	Leu	Gly	Gln	Pro	Ile	Phe
	210					215					220				
Pro	Gly	Asp	Ser	Gly	Val	Asp	Gln	Leu	Val	Glu	Ile	Ile	Lys	Val	Leu
225				230						235				240	
Gly	Thr	Pro	Thr	Arg	Glu	Gln	Ile	Arg	Glu	Met	Asn	Pro	Asn	Tyr	Thr
			245					250						255	
Glu	Phe	Lys	Phe	Pro	Gln	Ile	Lys	Ala	His	Pro	Trp	Thr	Lys	Val	Phe
			260					265					270		
Lys	Ser	Arg	Thr	Pro	Pro	Glu	Ala	Ile	Ala	Leu	Cys	Ser	Ser	Leu	Leu
		275				280						285			
Glu	Tyr	Thr	Pro	Ser	Ser	Arg	Leu	Ser	Pro	Leu	Glu	Ala	Cys	Ala	His
	290					295					300				
Ser	Phe	Phe	Asp	Glu	Leu	Arg	Cys	Leu	Gly	Thr	Gln	Leu	Pro	Asn	Asn
305				310					315					320	
Arg	Pro	Leu	Pro	Pro	Leu	Phe	Asn	Phe	Ser	Ala	Gly	Glu	Leu	Ser	Ile
			325					330						335	
Gln	Pro	Ser	Leu	Asn	Ala	Ile	Leu	Ile	Pro	Pro	His	Leu	Arg	Ser	
			340				345						350		

<210> 8

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> N-terminus addition sequence

<400> 8

Glu Phe Met Pro Thr Glu Ala Met Ala Ala Pro Lys Arg Val Ile



1 5 10 15

<210> 9  
<211> 9  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> N-terminus addition sequence

<400> 9  
Glu Tyr Met Pro Met Glu Gly Gly Gly  
1 5

<210> 10  
<211> 6  
<212> PRT  
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<220>  
<223> elution peptide

<400> 10  
Glu Tyr Met Pro Thr Asp  
1 5

<210> 11  
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<212> PRT  
<213> Artificial Sequence

<220>  
<223> Peptide substrate phosphorylatable by GSK3

<221> VARIANT  
<222> 2, 3, 4  
<223> Xaa = Any Amino Acid

<400> 11  
Ser Xaa Xaa Xaa Ser  
1 5